

OIPE

CRF Errors Corrected by the STIC System Branch

Serial Number: 09/847,208

CRF Processing Date: 5/16/2001

Edited by: [Signature]

Verified by: [Signature] (STIC staff)

**ENTERED**

- ☐ Changed a file from non-ASCII to ASCII
- ☐ Changed the margins in cases where the sequence text was "wrapped" down to the next line.
- ☐ Edited a format error in the Current Application Data section, specifically: \_\_\_\_\_
- ☐ Edited the Current Application Data section with the actual current number. The number inputted by the applicant was ☐ the prior application data; or ☐ other \_\_\_\_\_
- ☐ Added the mandatory heading and subheadings for "Current Application Data".
- ☐ Edited the "Number of Sequences" field. The applicant spelled out a number instead of using an integer.
- ☐ Changed the spelling of a mandatory field (the headings or subheadings), specifically: \_\_\_\_\_
- ☐ Corrected the SEQ ID NO when obviously incorrect. The sequence numbers that were edited were: \_\_\_\_\_
- ☐ Inserted or corrected a nucleic number at the end of a nucleic line. SEQ ID NO's edited: \_\_\_\_\_
- ☐ Corrected subheading placement. All responses must be on the same line as each subheading. If the applicant placed a response below the subheading, this was moved to its appropriate place.
- ☐ Inserted colons after headings/subheadings. Headings edited included: \_\_\_\_\_
- ☐ Deleted extra, invalid, headings used by an applicant, specifically: \_\_\_\_\_
- ☒ Deleted: ☒ non-ASCII "garbage" at the beginning/end of files; ☐ secretary initials/filename at end of file; ☐ page numbers throughout text; ☐ other invalid text, such as \_\_\_\_\_
- ☐ Inserted mandatory headings, specifically: \_\_\_\_\_
- ☐ Corrected an obvious error in the response, specifically: \_\_\_\_\_
- ☐ Edited identifiers where upper case is used but lower case is required, or vice versa.
- ☐ Corrected an error in the Number of Sequences field, specifically: \_\_\_\_\_
- ☐ A "Hard Page Break" code was inserted by the applicant. All occurrences had to be deleted.
- ☐ Deleted *ending* stop codon in amino acid sequences and adjusted the "(A)Length:" field accordingly (error due to a PatentIn bug). Sequences corrected: \_\_\_\_\_
- ☐ Other: \_\_\_\_\_

\*Examiner: The above corrections must be communicated to the applicant in the first Office Action. DO NOT send a copy of this form. 3/1/95

OIPE

RAW SEQUENCE LISTING  
 PATENT APPLICATION: US/09/847,208

DATE: 05/16/2001  
 TIME: 15:34:47

Input Set : A:\DES.txt  
 Output Set: N:\CRF3\05162001\I847208.raw

Does Not Comply  
 Corrected Diskette Needed

3 <110> APPLICANT: Saxon, Andrew  
 4       Zhang, Ke  
 5       Zhu, Daocheng  
 7 <120> TITLE OF INVENTION: FUSION MOLECULES AND TREATMENT OF  
 8       IgE-MEDIATED ALLERGIC DISEASES  
 10 <130> FILE REFERENCE: UC67.002A  
 C--> 12 <140> CURRENT APPLICATION NUMBER: US/09/847,208  
 C--> 12 <141> CURRENT FILING DATE: 2001-05-01  
 12 <160> NUMBER OF SEQ ID NOS: 177  
 14 <170> SOFTWARE: FastSEQ for Windows Version 4.0

# ERRORED SEQUENCES

6232 <210> SEQ ID NO: 177  
 6233 <211> LENGTH: 32  
 6234 <212> TYPE: PRT  
 6235 <213> ORGANISM: Homo sapiens  
 6237 <220> FEATURE:  
 6238 <221> NAME/KEY: UNSURE  
 6239 <222> LOCATION: 1, 4, 8, 10, 11, 12, 27, 30  
 6240 <223> OTHER INFORMATION: Xaa = any amino acid  
 6242 <400> SEQUENCE: 177  
 W--> 6243 Xaa Gln Gln Xaa Glu Leu Gln Xaa Asp Xaa Xaa Xaa Gln Ser Gln Leu  
       6244   1                  5                  10                  15  
 W--> 6245 Glu Arg Ala Asp Leu Arg Pro Gly Glu Gln Xaa Leu Met Xaa Lys Ile  
       6246           20                  25                  30  
 E--> 6251 W:\DOCS\BGY\SEQUENCE LISTING\UC67.003A.doc  
 E--> 6252 5  
 E--> 6254 1

VERIFICATION SUMMARY  
PATENT APPLICATION: US/09/847,208

DATE: 05/16/2001  
TIME: 15:34:49

Input Set : A:\DES.txt  
Output Set: N:\CRF3\05162001\I847208.raw

L:12 M:270 C: Current Application Number differs, Replaced Current Application No  
L:12 M:271 C: Current Filing Date differs, Replaced Current Filing Date  
L:3086 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:82  
L:3129 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:84  
L:3466 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:93  
L:3468 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:93  
L:3483 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:94  
L:4518 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:126  
L:6226 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:176  
L:6228 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:176  
L:6243 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:177  
L:6245 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:177  
L:6251 M:333 E: Wrong sequence grouping, Amino acids not in groups!  
L:6251 M:320 E: (1) Wrong Nucleic Acid Designator, NUMBER OF INVALID KEYS:2  
L:6252 M:332 E: (32) Invalid/Missing Amino Acid Numbering, SEQ ID:177  
M:332 Repeated in SeqNo=177  
L:6254 M:252 E: No. of Seq. differs, <211>LENGTH:Input:32 Found:34 SEQ:177